

GenCore version 5.1.6  
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OM nucleic - nucleic search, using sw model

Run on: May 4, 2005, 12:06:08 ; Search time 3110.34  
Seconds

(without alignments)  
5092.486 Million cell

updates/sec

Title: US-09-300-482-298  
Perfect score: 388  
Sequence: 1  
ggagaaagaaagaaaagatg.....ttaagcctgggacccccgt 388

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 45554873 seqs, 20411521753 residues

Total number of hits satisfying chosen parameters: 91109746

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Pending\_Patents\_NA\_Main:\*  
1: /cgn2\_6/ptodata/1/pna/PCTUS1\_COMB.seq:\*  
2: /cgn2\_6/ptodata/1/pna/PCTUS2\_COMB.seq:\*  
3: /cgn2\_6/ptodata/1/pna/PCTUS\_COMB.seq:\*  
4: /cgn2\_6/ptodata/1/pna/US06\_COMB.seq:\*  
5: /cgn2\_6/ptodata/1/pna/US07\_COMB.seq:\*  
6: /cgn2\_6/ptodata/1/pna/US080\_COMB.seq:\*  
7: /cgn2\_6/ptodata/1/pna/US081\_COMB.seq:\*  
8: /cgn2\_6/ptodata/1/pna/US082\_COMB.seq:\*  
9: /cgn2\_6/ptodata/1/pna/US083\_COMB.seq:\*  
10: /cgn2\_6/ptodata/1/pna/US084\_COMB.seq:\*  
11: /cgn2\_6/ptodata/1/pna/US085\_COMB.seq:\*  
12: /cgn2\_6/ptodata/1/pna/US086\_COMB.seq:\*  
13: /cgn2\_6/ptodata/1/pna/US087\_COMB.seq:\*